



OIIPE

RAW SEQUENCE LISTING

DATE: 01/23/2002

PATENT APPLICATION: US/10/038,694

TIME: 19:03:07

Input Set : A:\07083.0008U5.SEQ.TXT

Output Set: N:\CRF3\01232002\J038694.raw

ENTERED

p.5

4 <110> APPLICANT: Dixon, Eric
 5 Hutchins, Jeff T.
 6 Kuettner, Klaus E.
 7 Schmid, Thomas M.
 8 Schumacher, Barbara L.
 9 Su, Jui-Lan
 12 <120> TITLE OF INVENTION: SUPERFICIAL ZONE PROTEIN AND METHODS OF
 13 MAKING AND USING SAME
 16 <130> FILE REFERENCE: 07083.0008U5
 C--> 18 <140> CURRENT APPLICATION NUMBER: US/10/038,694
 C--> 18 <141> CURRENT FILING DATE: 2001-12-31
 18 <150> PRIOR APPLICATION NUMBER: 60/258,920
 19 <151> PRIOR FILING DATE: 2000-12-29
 21 <160> NUMBER OF SEQ ID NOS: 11
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 6
 27 <212> TYPE: PRT
 28 <213> ORGANISM: Artificial Sequence
 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
 32 synthetic construct
 34 <400> SEQUENCE: 1
 35 Asp Glu Ala Gly Ser Gly
 36 1 5
 38 <210> SEQ ID NO: 2
 39 <211> LENGTH: 188
 40 <212> TYPE: PRT
 41 <213> ORGANISM: Artificial Sequence
 43 <220> FEATURE:
 44 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
 45 synthetic construct
 47 <400> SEQUENCE: 2
 48 Met Ala Trp Lys Thr Leu Pro Ile Tyr Leu Leu Leu Leu Ser Val
 49 1 5 10 15
 50 Phe Val Ile Gln Gln Val Ser Ser Gln Asp Leu Ser Ser Cys Ala Gly
 51 20 25 30
 52 Arg Cys Gly Glu Gly Tyr Ser Arg Asp Ala Thr Cys Asn Cys Asp Tyr
 53 35 40 45
 54 Asn Cys Gln His Tyr Met Glu Cys Cys Pro Asp Phe Lys Arg Val Cys
 55 50 55 60
 56 Thr Ala Glu Leu Ser Cys Lys Gly Arg Cys Phe Glu Ser Phe Glu Arg
 57 65 70 75 80

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```

58 Gly Arg Glu Cys Asp Cys Asp Ala Gln Cys Lys Lys Tyr Asp Lys Cys
59           85           90           95
60 Cys Pro Asp Tyr Glu Ser Phe Cys Ala Glu Val Lys Asp Asn Lys Lys
61           100          105          110
62 Asn Arg Thr Lys Lys Lys Pro Thr Pro Lys Pro Pro Val Val Asp Glu
63           115           120           125
64 Ala Gly Ser Gly Leu Asp Asn Gly Asp Phe Lys Val Thr Thr Pro Asp
65           130           135           140
66 Thr Ser Thr Thr Gln His Asn Lys Val Ser Thr Ser Pro Lys Ile Thr
67 145           150           155           160
68 Thr Ala Lys Pro Ile Asn Pro Arg Pro Gln Ser Ser Pro Asn Ser Asp
69           165           170           175
70 Thr Ser Lys Glu Thr Ser Leu Thr Val Asn Lys Glu
71           180           185
73 <210> SEQ ID NO: 3
74 <211> LENGTH: 538
75 <212> TYPE: PRT
76 <213> ORGANISM: Artificial Sequence
78 <220> FEATURE:
79 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
80     synthetic construct
82 <400> SEQUENCE: 3
83 Pro Thr Thr Ile His Lys Ser Pro Asp Glu Ser Thr Pro Glu Leu Ser
84 1           5           10           15
85 Ala Glu Pro Thr Pro Lys Ala Leu Glu Asn Ser Pro Lys Glu Pro Gly
86           20           25           30
87 Val Pro Thr Thr Lys Thr Pro Ala Ala Thr Lys Pro Glu Met Thr Thr
88           35           40           45
89 Thr Ala Lys Asp Lys Thr Thr Glu Arg Asp Leu Arg Thr Thr Pro Glu
90           50           55           60
91 Thr Thr Thr Ala Ala Pro Lys Met Thr Lys Glu Thr Ala Thr Thr Thr
92 65           70           75           80
93 Glu Lys Thr Thr Glu Ser Lys Ile Thr Ala Thr Thr Thr Gln Val Thr
94           85           90           95
95 Ser Thr Thr Thr Gln Asp Thr Thr Pro Phe Lys Ile Thr Thr Leu Lys
96           100          105          110
97 Thr Thr Leu Ala Pro Lys Val Thr Thr Lys Lys Thr Ile Thr Thr
98           115          120          125
99 Thr Glu Ile Met Asn Lys Pro Glu Glu Thr Ala Lys Pro Lys Asp Arg
100          130          135          140
101 Ala Thr Asn Ser Lys Ala Thr Thr Pro Lys Pro Gln Lys Pro Thr Lys
102 145          150          155          160
103 Ala Pro Lys Lys Pro Thr Ser Thr Lys Lys Pro Lys Thr Met Pro Arg
104           165          170          175
105 Val Arg Lys Pro Lys Thr Thr Pro Thr Pro Arg Lys Met Thr Ser Thr
106           180          185          190
107 Met Pro Glu Leu Asn Pro Thr Ser Arg Ile Ala Glu Ala Met Leu Gln
108           195          200          205
109 Thr Thr Thr Arg Pro Asn Gln Thr Pro Asn Ser Lys Leu Val Glu Val

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```

110      210      215      220
111 Asn Pro Lys Ser Glu Asp Ala Gly Gly Ala Glu Gly Glu Thr Pro His
112 225      230      235      240
113 Met Leu Leu Arg Pro His Val Phe Met Pro Glu Val Thr Pro Asp Met
114      245      250      255
115 Asp Tyr Leu Pro Arg Val Pro Asn Gln Gly Ile Ile Ile Asn Pro Met
116      260      265      270
117 Leu Ser Asp Glu Thr Asn Ile Cys Asn Gly Lys Pro Val Asp Gly Leu
118      275      280      285
119 Thr Thr Leu Arg Asn Gly Thr Leu Val Ala Phe Arg Gly His Tyr Phe
120      290      295      300
121 Trp Met Leu Ser Pro Phe Ser Pro Pro Ser Pro Ala Arg Arg Ile Thr
122 305      310      315      320
123 Glu Val Trp Gly Ile Pro Ser Pro Ile Asp Thr Val Phe Thr Arg Cys
124      325      330      335
125 Asn Cys Glu Gly Lys Thr Phe Phe Phe Lys Asp Ser Gln Tyr Trp Arg
126      340      345      350
127 Phe Thr Asn Asp Ile Lys Asp Ala Gly Tyr Pro Lys Pro Ile Phe Lys
128      355      360      365
129 Gly Phe Gly Gly Leu Thr Gly Gln Ile Val Ala Ala Leu Ser Thr Ala
130      370      375      380
131 Lys Tyr Lys Asn Trp Pro Glu Ser Val Tyr Phe Phe Lys Arg Gly Gly
132 385      390      395      400
133 Ser Ile Gln Gln Tyr Ile Tyr Lys Gln Glu Pro Val Gln Lys Cys Pro
134      405      410      415
135 Gly Arg Arg Pro Ala Leu Asn Tyr Pro Val Tyr Gly Glu Met Thr Gln
136      420      425      430
137 Val Arg Arg Arg Arg Phe Glu Arg Ala Ile Gly Pro Ser Gln Thr His
138      435      440      445
139 Thr Ile Arg Ile Gln Tyr Ser Pro Ala Arg Leu Ala Tyr Gln Asp Lys
140      450      455      460
141 Gly Val Leu His Asn Glu Val Lys Val Ser Ile Leu Trp Arg Gly Leu
142 465      470      475      480
143 Pro Asn Val Val Thr Ser Ala Ile Ser Leu Pro Asn Ile Arg Lys Pro
144      485      490      495
145 Asp Gly Tyr Asp Tyr Tyr Ala Phe Ser Lys Asp Gln Tyr Tyr Asn Ile
146      500      505      510
147 Asp Val Pro Ser Arg Thr Ala Arg Ala Ile Thr Thr Arg Ser Gly Gln
148      515      520      525
149 Thr Leu Ser Lys Val Trp Tyr Asn Cys Pro
150      530      535

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152 <210> SEQ ID NO: 4

153 <211> LENGTH: 3

154 <212> TYPE: PRT

155 <213> ORGANISM: Artificial Sequence

157 <220> FEATURE:

158 <223> OTHER INFORMATION: Description of Artificial Sequence; note =

159 synthetic construct

161 <221> NAME/KEY: VARIANT

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Input Set : A:\07083.0008U5.SEQ.TXT

Output Set: N:\CRF3\01232002\J038694.raw

162 <222> LOCATION: 2
 163 <223> OTHER INFORMATION: Xaa is any amino acid except Pro
 165 <221> NAME/KEY: VARIANT
 166 <222> LOCATION: 3
 167 <223> OTHER INFORMATION: Xaa is either Thr or Ser
 169 <400> SEQUENCE: 4
 170 Asn Xaa Xaa
 171 1
 173 <210> SEQ ID NO: 5
 174 <211> LENGTH: 488
 175 <212> TYPE: DNA
 176 <213> ORGANISM: Artificial Sequence
 178 <220> FEATURE:
 179 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
 180 synthetic construct
 182 <400> SEQUENCE: 5
 183 atgcatgaaa cattocatta ctgtgttctg tgttgtttct gatcacaatt tatccaaatt 60
 184 atcagcgtga ggagagtggg agggatttag gatccactga acgtgttaaa cgtcacatac 120
 185 tgggtgtgcc tgtttaagga gctgactcgg gcttcgtaa ggcgcgcttg atcctcggag 180
 186 gggggggtgg acgcgcgcca agtagaatat acagtgtgtc cgttagaggt ttctgtgcag 240
 187 aagtaaaaga taacaagaag aacagaacta aaaagaaacc tacccecaaa ccaccagttg 300
 188 tagatgaagc tggaagtgga ttggacaatg gtgacttcaa ggtcacaact cctgacacgt 360
 189 ctaccaccca acacaataaa gtcagcacat ctccaagat cacaacagca aaaccaataa 420
 190 atcccagacc ccagtcttca cctaattctg ataactctaa agagacgtct ttgacagtga 480
 191 ataaagag 488
 193 <210> SEQ ID NO: 6
 194 <211> LENGTH: 1620
 195 <212> TYPE: DNA
 196 <213> ORGANISM: Artificial Sequence
 198 <220> FEATURE:
 199 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
 200 synthetic construct
 202 <400> SEQUENCE: 6
 203 cctaccacta tccacaaaag ccctgatgaa tcaactcctg agctttctgc agaaccacaca 60
 204 ccaaaagctc ttgaaaacag tcccaaggaa cctggtgtac ctacaactaa gactcctgca 120
 205 gcgactaaac ctgaaatgac tacaacagct aaagacaaga caacagaaag agacttacgt 180
 206 actacacctg aaactacaac tgctgcacct aagatgacaa aagagacagc aactacaaca 240
 207 gaaaaaacta ccgaatccaa aataacagct acaaccacac aagtaacatc taccacaact 300
 208 caagatacca caccattcaa aattactact cttaaaacaa ctactcttgc acccaaagta 360
 209 actacaacaa aaaagacaat tactaccact gagattatga acaaacctga agaaacagct 420
 210 aaaccaaaag acagagctac taattctaaa gcgacaactc ctaaacctca aaagccaacc 480
 211 aaagcaccca aaaaacccac ttctaccaaa aagccaaaaa caatgcctag agtgagaaaa 540
 212 ccaaagacga caccaactcc ccgcaagatg acatcaacaa tgccagaatt gaaccctacc 600
 213 tcaagaatag cagaagccat gctccaaacc accaccagac ctaaccaaac tccaaactcc 660
 214 aaactagttg aagtaaattcc aaagagtga gatgcaggtg gtgctgaagg agaaacacct 720
 215 catatgcttc tcaggcccca tgtgttcatg cctgaagtta ctcccagacat ggattactta 780
 216 ccgagagtac ccaatcaagg cattatcatc aatcccatgc tttccgatga gaccaatata 840
 217 tgcaatggta agccagtaga tggactgact actttgcgca atgggacatt agttgcattc 900
 218 cgaggtcatt atttctggat gctaagtcca ttcagtccac catctccagc tcgcagaatt 960

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DATE: 01/23/2002

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TIME: 19:03:07

Input Set : A:\07083.0008U5.SEQ.TXT

Output Set: N:\CRF3\01232002\J038694.raw

```

219 actgaagttt ggggtattcc ttccccatt gatactgttt ttactaggtg caactgtgaa 1020
220 ggaaaaactt tcttctttaa ggattctcag tactggcggt ttaccaatga tataaaagat 1080
221 gcagggtacc ccaaaccaat tttcaaagga tttggaggac taactggaca aatagtggca 1140
222 gcgctttcaa cagctaaata taagaactgg cctgaatctg tgtatttttt caagagaggt 1200
223 ggcagcattc agcagtatat ttataaacag gaacctgtac agaagtgcc tggaagaagg 1260
224 cctgctctaa attatccagt gtatggagaa atgacacagg ttaggagacg tcgctttgaa 1320
225 cgtgctatag gaccttctca aacacacacc atcagaattc aatattcacc tgccagactg 1380
226 gcttatcaag acaaagggtg ccttcataat gaagttaaag tgagtatact gtggagagga 1440
227 cttccaaatg tggttacctc agctatatca ctgcccaaca tcagaaaacc tgacggctat 1500
228 gattactatg ccttttctaa agatcaatac tataacattg atgtgcctag tagaacagca 1560
229 agagcaatta ctactcgttc tgggcagacc ttatccaaag tctggtacaa ctgtccttag 1620
231 <210> SEQ ID NO: 7
232 <211> LENGTH: 24
233 <212> TYPE: DNA
234 <213> ORGANISM: Artificial Sequence
236 <220> FEATURE:
237 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
238     synthetic construct
240 <400> SEQUENCE: 7
241 atggcatgga aaacacttcc catt                                     24
243 <210> SEQ ID NO: 8
244 <211> LENGTH: 24
245 <212> TYPE: DNA
246 <213> ORGANISM: Artificial Sequence
248 <220> FEATURE:
249 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
250     synthetic construct
252 <400> SEQUENCE: 8
253 ctaaggacag ttgtaccaga cttt                                     24
255 <210> SEQ ID NO: 9
256 <211> LENGTH: 4
257 <212> TYPE: PRT
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
262     synthetic construct
264 <400> SEQUENCE: 9
265 Phe Ala Cys Glu
266 1
268 <210> SEQ ID NO: 10
269 <211> LENGTH: 8
270 <212> TYPE: PRT
271 <213> ORGANISM: Artificial Sequence
273 <220> FEATURE:
274 <223> OTHER INFORMATION: Description of Artificial Sequence; note =
275     synthetic construct
277 <400> SEQUENCE: 10
278 Val Lys Asp Asn Lys Lys Asn Arg
279 1                                     5

```

Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/038,694

DATE: 01/23/2002

TIME: 19:03:08

Input Set : A:\07083.0008U5.SEQ.TXT

Output Set: N:\CRF3\01232002\J038694.raw

L:18 M:270 C: Current Application Number differs, Replaced Current Application No

L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

L:295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11